Appl. No. 09/700,712

Amendment dated: April 11, 2005 Reply to OA of: September 23, 2004

This listing of claims will replace all prior versions and listings of claims in the application.

**Listing of Claims**:

Claims 1-16(cancelled).

17(currently amended). A  $\Delta$  thy A strain of Vibrio cholerae deprived of its thy A gene in the chromosome and thus lacing lacking the functionality of the thy A gene comprising at least one episomal autonomously replicating DNA elements element having a functional thy A gene that enables the strain to grow in the absence of thymine in the growth medium and the one or several episomal autonomously replicating DNA elements further comprising a structural [[gen]] gene encoding a homologous or heterologous protein.

18(previously presented). A  $\Delta$  thy A strain of Vibrio cholerae wherein the strain has been deprived of its thy A gene by site-directed mutagenesis in the V. Cholerae chromosome for the deletion and/or insertion of nucleotides at the locus of the thy A gene.

19(currently amended). The  $\Delta$  thy A strain of Vibrio cholerae according to claim 17, wherein the <u>at least one</u> episomal autonomously replicating DNA element is a plasmid.

20(currently amended). The  $\Delta$  thy A strain of Vibrio cholerae according to claim 17, wherein the at least on [[ the]] episomal autonomously replicating DNA elements have element has a foreign thy A gene.

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21(previously presented) . The  $\Delta$  thy A strain of Vibrio cholerae according to claim 20, wherein the foreign thy A gene is an E. coli gene.

22(previously presented). The Δ *thy* A strain of *Vibrio cholerae* according to claim 17, wherein the encoded heterologous protein is selected from heat labile enterotoxin B-subunit of *Escherichia coli* (LTB) and *Schistosoma japonicum* glutathione S-transferase 26 kD protein (GST 26 kD).

23(previously presented). The  $\Delta$  *thy* A strain according to claim 17, wherein the *thy* A gene of the chromosome has the nucleotide sequence SEQ ID NO: 1, or said nucleotide sequence which has some natural or unnatural nucleotide extensions, truncations, deletions or additions that do not interfere with the natural function of the nucleotide sequence.